

## SEQUENCE LISTING

```
<110> GORING, Daphne R. et al.
 <120> PROLINE-RICH EXTENSIN-LIKE RECEPTOR KINASES
 <130> P 25,762-A USA
 <140> US 10/086,464
 <141> 2002-02-28
 <150> US 10/069,304
 <151> 2002-02-19
 <150> PCT/CA00/00966
 <151> 2000-08-18
 <150> US 60/149,466
 <151> 1999-08-19
<150> US 60/159,122
€151> 1999-10-13
<160> 27
<170> PatentIn Ver. 2.1
<210> 1
<211> 1944
<212> DNA
<213> Brassica napus
<220>
<221> CDS
<222> (1)..(1944)
<400> 1
ấtg tec teg geg ceg tet eeg ggg aet ggt teg eet eea tet eea eea
                                                                   48
Met Ser Ser Ala Pro Ser Pro Gly Thr Gly Ser Pro Pro Ser Pro Pro
  1
tca aac tcc aca acc act cct cct cca gct tcc gct cct ccc
                                                                   96
Ser Asn Ser Thr Thr Thr Pro Pro Pro Ala Ser Ala Pro Pro
             20
acc aca cct tct tct cct ccg ccg cca tcc act att ccg aca tct cct
                                                                   144
Thr Thr Pro Ser Ser Pro Pro Pro Pro Ser Thr Ile Pro Thr Ser Pro
         35
                             40
cct cct tct tct cgc tct aca cct tct gct cct cct cca tct cca cca
                                                                   192
Pro Pro Ser Ser Arg Ser Thr Pro Ser Ala Pro Pro Pro Ser Pro Pro
```

55

act Thr : 65	Pro	tct Ser	acg Thr	ccg Pro	gga Gly 70	tct Ser	cca Pro	cct Pro	cct Pro	ctt Leu 75	Pro	cag Gln	ccg Pro	tct Ser	cca Pro 80	240
ccc Pro	gct Ala	cca Pro	act Thr	acg Thr 85	Pro	gga Gly	tct Ser	cca Pro	ccc Pro 90	gca Ala	cct Pro	gtt Val	act Thr	cct Pro 95	cct Pro	288
act Thr	cga Arg	aac Asn	cct Pro 100	cca Pro	cct Pro	tca Ser	gtc Val	cca Pro 105	gga Gly	cca Pro	ccg Pro	tcc Ser	aat Asn 110	cct Pro	tca Ser	336
cgc Arg	gaa Glu	gga Gly 115	gga Gly	tct Ser	cct Pro	cga Arg	cct Pro 120	cca Pro	tct Ser	tct Ser	ccc Pro	tcg Ser 125	ccg Pro	ccg Pro	tct Ser	384
cct Pro	tct Ser 130	tcc Ser	gac Asp	ggt Gly	tta Leu	tca Ser 135	aca Thr	gga Gly	gtg Val	gtg Val	gtg Val 140	gga Gly	atc Ile	gcc Ala	atc Ile	432
gga Gly 145	gga Gly	gtc Val	gct Ala	ctg Leu	ctt Leu 150	gtg Val	ata Ile	gtg Val	act Thr	ctg Leu 155	att Ile	tgt Cys	ctc Leu	ctc Leu	tgt Cys 160	480
aag Lys	aag Lys	aaa Lys	cga Arg	cgg Arg 165	aga Arg	gac Asp	gaa Glu	gaa Glu	gat Asp 170	gct Ala	tac Tyr	tat Tyr	gtt Val	cct Pro 175	ccg Pro	528
cca Pro	cct Pro	cct Pro	cct Pro 180	ggt Gly	ccc Pro	aaa Lys	gcc Ala	gga Gly 185	gga Gly	cct Pro	tac Tyr	ggt Gly	gga Gly 190	cag Gln	cag Gln	576
caa Gln	caa Gln	tgg Trp 195	cgg Arg	caa Gln	caa Gln	aac Asn	gca Ala 200	aca Thr	cca Pro	ccg Pro	tca Ser	gat Asp 205	cat His	gtc Val	gtg Val	624
acg Thr	tca Ser 210	cta Leu	cca Pro	cca Pro	cca Pro	cct Pro 215	aag Lys	gct Ala	cca Pro	tct Ser	cca Pro 220	cca Pro	cgg Arg	caa Gln	cct Pro	672
cct Pro 225	cca Pro	cct Pro	cca Pro	cca Pro	ccg Pro 230	cct Pro	ttc Phe	atg Met	agc Ser	agc Ser 235	agc Ser	ggc Gly	ggc Gly	tcc Ser	gac Asp 240	720
tac Tyr	tcg Ser	gac Asp	Arg	cca Pro 245	gtt Val	ctt Leu	cct Pro	cca Pro	ccg Pro 250	tct Ser	cca Pro	Gly 999	ctt Leu	gtg Val 255	tta Leu	768
ggc Gly	ttc Phe	Ser	aaa Lys 260	agc Ser	act Thr	ttc Phe	aca Thr	tac Tyr 265	gag Glu	gag Glu	cta Leu	gct Ala	aga Arg 270	gcc Ala	acc Thr	816
aat Āsn	ggt Gly	ttc Phe	tcc Ser	gag Glu	gcg Ala	aac Asn	ttg Leu	tta Leu	gga Gly	caa Gln	ggc Gly	Gly 999	ttc Phe	ggt Gly	tac Tyr	864

•

gtg Val	cac His 290	Lys	ggt Gly	gtg Val	ttg Leu	cct Pro 295	agt Ser	Gly 999	aaa Lys	gaa Glu	gtt Val 300	Ala	gtg Val	aag Lys	cag Gln	912
ttg Ļeu 305	aaa Lys	gtt Val	Gly aaa	agt Ser	ggt Gly 310	cag Gln	gga Gly	gag Glu	agg Arg	gag Glu 315	ttt Phe	cag Gln	gca Ala	gag Glu	gtt Val 320	960
gag Glu	atc Ile	atc Ile	agc Ser	aga Arg 325	gtt Val	cac His	cac His	agg Arg	cat His 330	ctg Leu	gtg Val	tct Ser	ctt Leu	gtt Val 335	ggt Gly	1008
tat Tyr	tgc Cys	atc Ile	gcc Ala 340	ggt Gly	gcc Ala	aaa Lys	aga Arg	ttg Leu 345	ctt Leu	gtc Val	tat Tyr	gag Glu	ttt Phe 350	gtt Val	cct Pro	1056
aac Asn	aac Asn	aat Asn 355	ctc Leu	gag Glu	ctt Leu	cac His	ctc Leu 360	cat His	ggc Gly	gag Glu	gga Gly	cgg Arg 365	cct Pro	aca Thr	atg Met	1104
gaa Glu	tgg Trp 370	agc Ser	acc Thr	aga Arg	ttg Leu	aag Lys 375	att Ile	gct Ala	ctt Leu	gga Gly	tct Ser 380	gct Ala	aaa Lys	gga Gly	ctt Leu	1152
tct Ser 385	tat Tyr	ctt Leu	cat His	gaa Glu	gat Asp 390	tgc Cys	aat Asn	cct Pro	aaa Lys	atc Ile 395	att Ile	cac His	cgt Arg	gat Asp	atc Ile 400	1200
aag Lys	gct Ala	tca Ser	aac Asn	ata Ile 405	ttg Leu	ata Ile	gat Asp	ttc Phe	aag Lys 410	ttt Phe	gaa Glu	gct Ala	aag Lys	gtt Val 415	gct Ala	1248
gat Asp	ttt Phe	ggt Gly	ctt Leu 420	gct Ala	aag Lys	att Ile	gct Ala	tct Ser 425	gat Asp	aca Thr	aac Asn	acg Thr	cat His 430	gta Val	tca Ser	1296
aca Thr	cgt Arg	gtg Val 435	atg Met	gga Gly	acc Thr	ttt Phe	999 Gly 440	tac Tyr	ttg Leu	gct Ala	ccg Pro	gaa Glu 445	tac Tyr	gct Ala	gca Ala	1344
agc	gga Gly 450	aag Lys	ctc Leu	acg Thr	gag Glu	aag Lys 455	tct Ser	gac Asp	gtt Val	ttc Phe	tca Ser 460	ttt Phe	ggc Gly	gtt Val	gtg Val	1392
ctt Leu 465	ttg Leu	gag Glu	ctc Leu	att Ile	act Thr 470	gga Gly	cgt Arg	cga Arg	ccc Pro	gtt Val 475	gat Asp	gcc Ala	aac Asn	aat Asn	gtc Val 480	1440
tat Tyr	gta Val	gat Asp	gac Asp	agc Ser 485	tta Leu	gtt Val	gac Asp	tgg Trp	gca Ala 490	cga Arg	cca Pro	ttg Leu	ctt Leu	aac Asn 495	cga Arg	1488

5																
gca Ala	tct Ser	gag Glu	caa Gln 500	gga Gly	gac Asp	ttt Phe	gag Glu	ggt Gly 505	tta Leu	gct Ala	gat Asp	gca Ala	aag Lys 510	atg Met	aat Asn	1536
aat Asn	Gly 999	tat Tyr 515	gac Asp	aga Arg	gag Glu	gag Glu	atg Met 520	gct Ala	cgc Arg	atg Met	gtt Val	gct Ala 525	tgt Cys	gct Ala	gcg Ala	1584
gct Ala	tgt Cys 530	gtt Val	cgc Arg	cat His	tca Ser	gct Ala 535	cgc Arg	cgc Arg	aga Arg	cct Pro	cgc Arg 540	atg Met	agc Ser	cag Gln	att Ile	1632
											gat Asp					1680
											tac Tyr					1728
											aag Lys					1776
atg Met	gca Ala	ctt Leu 595	gga Gly	act Thr	caa Gln	gag Glu	tac Tyr 600	aac Asn	gcc Ala	acg Thr	ggt Gly	gag Glu 605	tac Tyr	agt Ser	aat Asn	1824
											tca Ser 620					1872
caa Gln 625	acc Thr	aca Thr	cgc Arg	gaa Glu	atg Met 630	gag Glu	atg Met	Gly 999	aag Lys	att Ile 635	aag Lys	aga Arg	acc Thr	ggt Gly	cag Gln 640	1920
					tct Ser	ctt Leu	taa									1944

<210> 2

<211> 647

<212> PRT

<213> Brassica napus

<400> 2

Thr Pro Ser Thr Pro Gly Ser Pro Pro Pro Leu Pro Gln Pro Ser Pro Pro Ala Pro Thr Thr Pro Gly Ser Pro Pro Ala Pro Val Thr Pro Pro Thr Arg Asn Pro Pro Pro Ser Val Pro Gly Pro Pro Ser Asn Pro Ser Ärg Glu Gly Gly Ser Pro Arg Pro Pro Ser Ser Pro Ser Pro Pro Ser Pro Ser Ser Asp Gly Leu Ser Thr Gly Val Val Gly Ile Ala Ile Gly Gly Val Ala Leu Leu Val Ile Val Thr Leu Ile Cys Leu Leu Cys Lys Lys Lys Arg Arg Arg Asp Glu Glu Asp Ala Tyr Tyr Val Pro Pro Pro Pro Pro Gly Pro Lys Ala Gly Gly Pro Tyr Gly Gly Gln Gln Gln Gln Trp Arg Gln Gln Asn Ala Thr Pro Pro Ser Asp His Val Val Thr Ser Leu Pro Pro Pro Lys Ala Pro Ser Pro Pro Arg Gln Pro Pro Pro Pro Pro Pro Phe Met Ser Ser Gly Gly Ser Asp Tyr Ser Asp Arg Pro Val Leu Pro Pro Pro Ser Pro Gly Leu Val Leu Gly Phe Ser Lys Ser Thr Phe Thr Tyr Glu Glu Leu Ala Arg Ala Thr Asn Gly Phe Ser Glu Ala Asn Leu Leu Gly Gln Gly Phe Gly Tyr Val His Lys Gly Val Leu Pro Ser Gly Lys Glu Val Ala Val Lys Gln Leu Lys Val Gly Ser Gly Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Glu Ile Ile Ser Arg Val His His Arg His Leu Val Ser Leu Val Gly Tyr Cys Ile Ala Gly Ala Lys Arg Leu Leu Val Tyr Glu Phe Val Pro Asn Asn Asn Leu Glu Leu His Leu His Gly Glu Gly Arg Pro Thr Met Glu Trp Ser Thr Arg Leu Lys Ile Ala Leu Gly Ser Ala Lys Gly Leu Ser Tyr Leu His Glu Asp Cys Asn Pro Lys Ile Ile His Arg Asp Ile Lys Ala Ser Asn Ile Leu Ile Asp Phe Lys Phe Glu Ala Lys Val Ala Asp Phe Gly Leu Ala Lys Ile Ala Ser Asp Thr Asn Thr His Val Ser Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ala Ser Gly Lys Leu Thr Glu Lys Ser Asp Val Phe Ser Phe Gly Val Val Leu Leu Glu Leu Ile Thr Gly Arg Arg Pro Val Asp Ala Asn Asn Val 475. Tyr Val Asp Asp Ser Leu Val Asp Trp Ala Arg Pro Leu Leu Asn Arg 

Ala Ser Glu Gln Gly Asp Phe Glu Gly Leu Ala Asp Ala Lys Met Asn 505 Asn Gly Tyr Asp Arg Glu Glu Met Ala Arg Met Val Ala Cys Ala Ala 520 Ala Cys Val Arg His Ser Ala Arg Arg Arg Pro Arg Met Ser Gln Ile 535 540 Val Arg Ala Leu Glu Gly Asn Val Ser Leu Ser Asp Leu Asn Glu Gly 550 Met Arg Pro Gly Gln Ser Asn Val Tyr Ser Ser Tyr Gly Gly Ser Thr 570 Asp Tyr Asp Ser Ser Gln Tyr Asn Glu Asp Met Lys Lys Phe Arg Lys 580 585 Met Ala Leu Gly Thr Gln Glu Tyr Asn Ala Thr Gly Glu Tyr Ser Asn 600 Pro Thr Ser Asp Tyr Gly Leu Tyr Pro Ser Gly Ser Ser Ser Glu Gly 615 620 Gln Thr Thr Arg Glu Met Glu Met Gly Lys Ile Lys Arg Thr Gly Gln 635 Gly Tyr Ser Gly Pro Ser Leu <210> 3 <211> 2189 <212> DNA <213> Brassica napus ₹220> <221> CDS <222> (1)...(2189)<223> <400> ttaactctct ggtctccgtg tctcctctt tctcctgctg cttcctttta acactctctt 60 catttgcctt tttgatttag atccaaagaa gcagac atg tcc tcg gcg ccg tct 114 Met Ser Ser Ala Pro Ser ccg ggg act ggt tcg cct cca tct cca cca tca aac tcc aca acc 162 Pro Gly Thr Gly Ser Pro Pro Ser Pro Pro Ser Asn Ser Thr Thr 15 act cet eet eea get tee get eet eet eee ace aca eet tet tet eet 210 Thr Pro Pro Pro Ala Ser Ala Pro Pro Pro Thr Thr Pro Ser Ser Pro 25 30 35 ccg ccg cca tcc act att ccg aca tct cct cct tct tct cgc tct 258 Pro Pro Pro Ser Thr Ile Pro Thr Ser Pro Pro Pro Ser Ser Arg Ser

								cca Pro								306
								tct Ser								354
								cct Pro 95								402
								cct Pro								450
								ccg Pro								498
								gcc Ala								546
								ctc Leu								594
								cct Pro 175								642
								cag Gln								690
								gtc Val								738
cct Pro 215	aag Lys	gct Ala	cca Pro	tct Ser	cca Pro 220	cca Pro	cgg Arg	caa Gln	cct Pro	cct Pro 225	cca Pro	cct Pro	cca Pro	cca Pro	ccg Pro 230	786
								tcc Ser								834
								gtg Val 255								882
								gcc Ala								930

.

aac Asr	ttg Leu 280	ı ьеи	gga Gly	caa Gln	ggc Gly	999 285	ttc Phe	ggt Gly	tac Tyr	gtg Val	cac His	Lys	ggt Gly	gtg Val	ttg Leu	978
cct Pro 295	Ser	. Gly aaa	aaa Lys	gaa Glu	gtt Val 300	gct Ala	gtg Val	aag Lys	cag Gln	ttg Leu 305	Lys	gtt Val	Gly 999	agt Ser	ggt Gly 310	1026
Çag Gln	gga Gly	gag Glu	agg Arg	gag Glu 315	ttt Phe	cag Gln	gca Ala	gag Glu	gtt Val 320	gag Glu	atc Ile	atc Ile	agc Ser	aga Arg 325	Val	1074
cac His	cac His	agg Arg	cat His 330	ctg Leu	gtg Val	tct Ser	ctt Leu	gtt Val 335	ggt Gly	tat Tyr	tgc Cys	atc Ile	gcc Ala 340	ggt Gly	gcc Ala	1122
aaa Lys	aga Arg	ttg Leu 345	ctt Leu	gtc Val	tat Tyr	gag Glu	ttt Phe 350	gtt Val	cct Pro	aac Asn	aac Asn	aat Asn 355	ctc Leu	gag Glu	ctt Leu	1170
cac His	ctc Leu 360	cat His	ggc Gly	gag Glu	gga Gly	cgg Arg 365	cct Pro	aca Thr	atg Met	gaa Glu	tgg Trp 370	agc Ser	acc Thr	aga Arg	ttg Leu	1218
āag Ļys 375	att Ile	gct Ala	ctt Leu	gga Gly	tct Ser 380	gct Ala	aaa Lys	gga Gly	ctt Leu	tct Ser 385	tat Tyr	ctt Leu	cat His	gaa Glu	gat Asp 390	1266
tgc Cys	aat Asn	cct Pro	aaa Lys	atc Ile 395	att Ile	cac His	cgt Arg	gat Asp	atc Ile 400	aag Lys	gct Ala	tca Ser	aac Asn	ata Ile 405	ttg Leu	1314
ata Ile	gat Asp	ttc Phe	aag Lys 410	ttt Phe	gaa Glu	gct Ala	aag Lys	gtt Val 415	gct Ala	gat Asp	ttt Phe	ggt Gly	ctt Leu 420	gct Ala	aag Lys	1362
att Ile	gct Ala	tct Ser 425	gat Asp	aca Thr	aac Asn	acg Thr	cat His 430	gta Val	tca Ser	aca Thr	cgt Arg	gtg Val 435	atg Met	gga Gly	acc Thr	1410
ttt Phe	999 Gly 440	tac Tyr	ttg Leu	gct Ala	ccg Pro	gaa Glu 445	tac Tyr	gct Ala	gca Ala	agc Ser	gga Gly 450	aag Lys	ctc Leu	acg Thr	gag Glu	1458
aag Lys 455	tct Ser	gac Asp	gtt Val	ttc Phe	tca Ser 460	ttt Phe	ggc Gly	gtt Val	gtg Val	ctt Leu 465	ttg Leu	gag Glu	ctc Leu	att Ile	act Thr 470	1506
gga Gly	cgt Arg	cga Arg	Pro	gtt Val 475	gat Asp	gcc Ala	aac Asn	Asn	gtc Val 480	tat Tyr	gta Val	gat Asp	gac Asp	agc Ser 485	tta Leu	1554

ģtt Val	gac Asp	tgg Trp	gca Ala 490	cga Arg	cca Pro	ttg Leu	ctt Leu	aac Asn 495	cga Arg	gca Ala	tct Ser	gag Glu	caa Gln 500	gga Gly	gac Asp	1602
ttt Phe	gag Glu	ggt Gly 505	tta Leu	gct Ala	gat Asp	gca Ala	aag Lys 510	atg Met	aat Asn	aat Asn	Gly 999	tat Tyr 515	gac Asp	aga Arg	gag Glu	1650
gag Glu	atg Met 520	gct Ala	cgc Arg	atg Met	gtt Val	gct Ala 525	tgt Cys	gct Ala	gcg Ala	gct Ala	tgt Cys 530	gtt Val	cgc Arg	cat His	tca Ser	1698
gct Ala 535	cgc Arg	cgc Arg	aga Arg	cct Pro	cgc Arg 540	atg Met	agc Ser	cag Gln	att Ile	gtg Val 545	cgt Arg	gcg Ala	tta Leu	gaa Glu	gga Gly 550	1746
aat Asn	gta Val	tca Ser	ctg Leu	tca Ser 555	gat Asp	ctt Leu	aac Asn	gaa Glu	999 Gly 560	atg Met	aga Arg	cca Pro	ggt Gly	caa Gln 565	agc Ser	1794
aat • Asn	gta Val	tac Tyr	agc Ser 570	tca Ser	tac Tyr	gga Gly	gga Gly	agc Ser 575	acc Thr	gat Asp	tat Tyr	gac Asp	tcg Ser 580	agc Ser	cag Gln <sub>,</sub>	1842
tac Tyr	aat Asn	gaa Glu 585	gac Asp	atg Met	aag Lys	aag Lys	ttt Phe 590	agg Arg	aaa Lys	atg Met	gca Ala	ctt Leu 595	gga Gly	act Thr	caa Gln	1890
gag Glu	tac Tyr 600	aac Asn	gcc Ala	acg Thr	ggt Gly	gag Glu 605	tac Tyr	agt Ser	aat Asn	ccg Pro	acc Thr 610	agt Ser	gac Asp	tat Tyr	gga Gly	1938
ctg Leu 615	tac Tyr	ccg Pro	tct Ser	ggt Gly	tca Ser 620	agc Ser	agc Ser	gag Glu	ggc Gly	caa Gln 625	acc Thr	aca Thr	cgc Arg	gaa Glu	atg Met 630	1986
gag Glu	atg Met	Gly aaa	aag Lys	att Ile 635	aag Lys	aga Arg	acc Thr	ggt Gly	cag Gln 640	ggt Gly	tat Tyr	agt Ser	gga Gly	cct Pro 645	tct Ser	2034
ctt Leu	t aa	acca	gatg.	gga	ıgaga	aat	tgaa	ıgggt	gt t	tttt	catt	a tt	tttt	taaa	ı	2088
acto	rtaaa	ga t	atga	gaaa	a tt	gcct	tact	cta	atta	aaa	ccac	tacg	at a	taag	gttat	2148
aata	cgtt	tt g	aatt	ggtt	t tt	aaaa	aaaa	aaa	aaaa	aaa	a					2189

\$210> 4
\$211> 647
\$212> PRT

<213> Brassica napus

Met Ser Ser Ala Pro Ser Pro Gly Thr Gly Ser Pro Pro Pro Pro Ser Asn Ser Thr Thr Thr Pro Pro Pro Ala Ser Ala Pro Pro Thr Thr Pro Ser Ser Pro Pro Pro Pro Ser Thr Ile Pro Thr Ser Pro Pro Pro Ser Ser Arg Ser Thr Pro Ser Ala Pro Pro Pro Ser Pro Pro Thr Pro Ser Thr Pro Gly Ser Pro Pro Pro Leu Pro Gln Pro Ser Pro Pro Ala Pro Thr Thr Pro Gly Ser Pro Pro Ala Pro Val Thr Pro Pro Thr Arg Asn Pro Pro Pro Ser Val Pro Gly Pro Pro Ser Asn Pro Ser Arg Glu Gly Ser Pro Arg Pro Pro Ser Ser Pro Ser Pro Pro Ser Pro Ser Ser Asp Gly Leu Ser Thr Gly Val Val Val Gly Ile Ala Ile Gly Gly Val Ala Leu Leu Val Ile Val Thr Leu Ile Cys Leu Leu Cys Lys Lys Lys Arg Arg Arg Asp Glu Glu Asp Ala Tyr Tyr Val Pro Pro Pro Pro Pro Gly Pro Lys Ala Gly Gly Pro Tyr Gly Gly Gln Gln Gln Gln Trp Arg Gln Gln Asn Ala Thr Pro Pro Ser Asp His Val Val Thr Ser Leu Pro Pro Pro Lys Ala Pro Ser Pro Pro Arg Gln Pro Pro Pro Pro Pro Pro Pro Phe Met Ser Ser Gly Gly Ser Asp Tyr Ser Asp Arg Pro Val Leu Pro Pro Pro Ser Pro Gly Leu Val Leu Gly Phe Ser Lys Ser Thr Phe Thr Tyr Glu Glu Leu Ala Arg Ala Thr Asn Gly Phe Ser Glu Ala Asn Leu Leu Gly Gln Gly Gly Phe Gly Tyr Val His Lys Gly Val Leu Pro Ser Gly Lys Glu Val Ala Val Lys Gln Leu Lys Val Gly Ser Gly Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Glu Ile Ile Ser Arg Val His His Arg His Leu Val Ser Leu Val Gly Tyr Cys Ile Ala Gly Ala Lys Arg Leu Leu Val Tyr Glu Phe Val Pro Asn Asn Asn Leu Glu Leu His Leu His Gly Glu Gly Arg Pro Thr Met Glu Trp Ser Thr Arg Leu Lys Ile Ala Leu Gly Ser Ala Lys Gly Leu Ser Tyr Leu His Glu Asp Cys Asn Pro Lys Ile Ile His Arg Asp Ile Lys Ala Ser Asn Ile Leu Ile Asp Phe Lys Phe Glu Ala Lys Val Ala Asp Phe Gly Leu Ala Lys Ile Ala Ser Asp Thr Asn Thr His Val Ser

420 425 430 Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ala 440 Ser Gly Lys Leu Thr Glu Lys Ser Asp Val Phe Ser Phe Gly Val Val 455 460 Leu Leu Glu Leu Ile Thr Gly Arg Pro Val Asp Ala Asn Asn Val 470 475 Tyr Val Asp Asp Ser Leu Val Asp Trp Ala Arg Pro Leu Leu Asn Arg 485 490 Ala Ser Glu Gln Gly Asp Phe Glu Gly Leu Ala Asp Ala Lys Met Asn 505 510 Asn Gly Tyr Asp Arg Glu Glu Met Ala Arg Met Val Ala Cys Ala Ala 520 Ala Cys Val Arg His Ser Ala Arg Arg Arg Pro Arg Met Ser Gln Ile 535 540 Val Arg Ala Leu Glu Gly Asn Val Ser Leu Ser Asp Leu Asn Glu Gly 550 555 Met Arg Pro Gly Gln Ser Asn Val Tyr Ser Ser Tyr Gly Gly Ser Thr 565 570 Asp Tyr Asp Ser Ser Gln Tyr Asn Glu Asp Met Lys Lys Phe Arg Lys 580 585 Met Ala Leu Gly Thr Gln Glu Tyr Asn Ala Thr Gly Glu Tyr Ser Asn Pro Thr Ser Asp Tyr Gly Leu Tyr Pro Ser Gly Ser Ser Ser Glu Gly 615 620 Gln Thr Thr Arg Glu Met Glu Met Gly Lys Ile Lys Arg Thr Gly Gln 630 635 Gly Tyr Ser Gly Pro Ser Leu 645

<210> 5
<211> 721
<212> PRT
<213> Brassica napus

## <400> 5

Leu Thr Leu Trp Ser Pro Cys Leu Leu Ser Ser Pro Ala Ala Ser Phe 1 5 10 15

His Ser Leu His Leu Pro Phe Phe Arg Ser Lys Glu Ala Asp Met Ser 20 25 30

Šer Ala Pro Ser Pro Gly Thr Gly Ser Pro Pro Ser Pro Pro Ser Asn 35 40 45

Ser Thr Thr Thr Pro Pro Pro Ala Ser Ala Pro Pro Pro Thr Thr 50 55 60

Pro Ser Ser Pro Pro Pro Pro Ser Thr Ile Pro Thr Ser Pro Pro Ser Ser Arg Ser Thr Pro Ser Ala Pro Pro Pro Ser Pro Pro Thr Pro Ser Thr Pro Gly Ser Pro Pro Pro Leu Pro Gln Pro Ser Pro Pro Ala Pro Thr Thr Pro Gly Ser Pro Pro Ala Pro Val Thr Pro Pro Thr Arg Asn Pro Pro Pro Ser Val Pro Gly Pro Pro Ser Asn Pro Ser Arg Glu Gly Gly Ser Pro Arg Pro Pro Ser Ser Pro Ser Pro Pro Ser Pro Ser Ser Asp Gly Leu Ser Thr Gly Val Val Gly Ile Ala Ile Gly Gly Val Ala Leu Leu Val Ile Val Thr Leu Ile Cys Leu Leu Cys Lys Lys Arg Arg Arg Asp Glu Glu Asp Ala Tyr Tyr Val Pro Pro Pro Pro Pro Gly Pro Lys Ala Gly Gly Pro Tyr Gly Gly Gln Gln Gln Trp Arg Gln Gln Asn Ala Thr Pro Pro Ser Asp His Val Val Thr Ser Leu Pro Pro Pro Lys Ala Pro Ser Pro Pro Arq Gln Pro Pro Pro Pro Pro Pro Phe Met Ser Ser Ser Gly Gly Ser Asp Tyr Ser - 270

Asp Arg Pro Val Leu Pro Pro Pro Ser Pro Gly Leu Val Leu Gly Phe

## 275 280 285

Ser Lys Ser Thr Phe Thr Tyr Glu Glu Leu Ala Arg Ala Thr Asn Gly Phe Ser Glu Ala Asn Leu Leu Gly Gln Gly Gly Phe Gly Tyr Val His Lys Gly Val Leu Pro Ser Gly Lys Glu Val Ala Val Lys Gln Leu Lys Val Gly Ser Gly Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Glu Ile Ile Ser Arg Val His His Arg His Leu Val Ser Leu Val Gly Tyr Cys Ile Ala Gly Ala Lys Arg Leu Leu Val Tyr Glu Phe Val Pro Asn Asn Äsn Leu Glu Leu His Leu His Gly Glu Gly Arg Pro Thr Met Glu Trp Ser Thr Arg Leu Lys Ile Ala Leu Gly Ser Ala Lys Gly Leu Ser Tyr Leu His Glu Asp Cys Asn Pro Lys Ile Ile His Arg Asp Ile Lys Ala Ser Asn Ile Leu Ile Asp Phe Lys Phe Glu Ala Lys Val Ala Asp Phe Gly Leu Ala Lys Ile Ala Ser Asp Thr Asn Thr His Val Ser Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ala Ser Gly Lys Leu Thr Glu Lys Ser Asp Val Phe Ser Phe Gly Val Val Leu Leu 

Glu Leu Ile Thr Gly Arg Arg Pro Val Asp Ala Asn Asn Val Tyr Val 500 505 510

Asp Asp Ser Leu Val Asp Trp Ala Arg Pro Leu Leu Asn Arg Ala Ser 515 520 525

Glu Gln Gly Asp Phe Glu Gly Leu Ala Asp Ala Lys Met Asn Asn Gly 530 540

Tyr Asp Arg Glu Glu Met Ala Arg Met Val Ala Cys Ala Ala Ala Cys 545 550 555 560

Val Arg His Ser Ala Arg Arg Pro Arg Met Ser Gln Ile Val Arg 565 570 575

Ala Leu Glu Gly Asn Val Ser Leu Ser Asp Leu Asn Glu Gly Met Arg 580 585 590

Pro Gly Gln Ser Asn Val Tyr Ser Ser Tyr Gly Gly Ser Thr Asp Tyr 595 600 605

Asp Ser Ser Gln Tyr Asn Glu Asp Met Lys Lys Phe Arg Lys Met Ala 610 620

Leu Gly Thr Gln Glu Tyr Asn Ala Thr Gly Glu Tyr Ser Asn Pro Thr 625 630 635 640

Ser Asp Tyr Gly Leu Tyr Pro Ser Gly Ser Ser Ser Glu Gly Gln Thr 645 650 655

Thr Arg Glu Met Glu Met Gly Lys Ile Lys Arg Thr Gly Gln Gly Tyr 660 665 670

Ser Gly Pro Ser Leu Thr Arg Trp Glu Arg Asn Arg Val Phe Phe His 675 680 685

Tyr Phe Phe Lys Thr Val Lys Ile Glu Asn Cys Leu Thr Leu Ile Lys 690 695 700

Thr Thr Thr Ile Gly Tyr Asn Thr Phe Ile Gly Phe Lys Lys Lys

705 710 715 720

Lys

<210> 6
<211> 2231
<212> DNA
<213> Arabidopsis thaliana

<400> 6 tagaaaaaaa aaaatgtcag acttaggcga gtcgccgagt tcttcaccac cagcaccacc 60 agetgatace getectecae cagagaetee ateagaaaae teagetette caectgttga 120 ttcctctcct cctagtccac cagctgattc atcatcaaca ccgccgctgt cagaaccatc 180 cactectect ccagatteac agetteetee tttacetteg attetteete egetaacaga 240 ttetecacet ceaectteeg attettetee accegttgat teaacceett eteegeegee 300 accgacgtca aacgaatctc cttctcctcc agaagattcc gaaacaccac ctgctccacc 360 aaatgaatcc aatgacaaca accetectee gteteaagat etteaatege eteeteeate 420 gtcqccgtcg ccgaatgtag gacccacaaa cccgqaatca ccaccgttac aatctcctcc 480 agetecacea geateagate etacaaatte acegeeaget teaceattag aceetaceaa 540 tectececca atacaaccat caggaccage caettetect ceggetaate ecaaegetee 600 geegageeca tteeceacag taccacecaa aacteettet agtggaeetg tggtgtetee 660 Steteteaca teceetagta aaggaaetee taeteeaaae caaggeaatg gagatggegg 720 tggcggtggt ggcggctatc aagggaagac tatggttggt atggctgtag ccggtttcgc 780 åatcatggcg cttataggcg ttgtgttctt agtgagaaga aagaaaaaga gaaacattga 840 tagctataat cactcacagt acttgccaca tcccaatttc tctgttaaat cagatggatt 900 cttatacggt caagatccag gtaaaggata ctcctctggt cctaatggtt caatgtataa 960 caattcacag caacaacaat cctctatggg aaacagttat ggtacagctg gtggtggtta 1020 tecteateat caaatgeaat caagtggeac acetgaetet getataeteg gaagtggeea 1080 gactcattte agttacgaag agcttgetga gataacacaa ggctttgete qeaaaaacat 1140 tcttggagaa ggcggatttg gatgtgtcta taaaggtaca ttgcaggatg gtaaagttgt 1200 tgcggttaag cagcttaaag ctggaagtgg acaaggtgac cgtgaattca aagcagaggt 1260 tgagatcatc agccgcgttc atcatcgcca tttggtctct ctggttggtt actgcatttc 1320 agaccagcat agattgctta tctatgagta tgtttctaat caaaccttgg agcatcattt 1380 gcatgagtgg tctaagagag tccggatcgc tataggatca gccaaagggt tggcatatct 1440 tcacqaaqac tqtcatccqa aaatcattca caqaqatata aaqtcaqcaa atattcttct 1500 agatgatgaa tatgaagctc aggcaataat gaaatcctcc ttttcgttaa atctatctta 1560 tgactgtaaa gttttagttg ctgattttgg acttgctaga ctcaatgata caacacaaac 1620 tcatgtttca actcgggtta tgggaacctt cgggtaccta gcgccggaat atgcatcaag 1680 Eggaaaattg actgatagat ccgatgtatt ctcattcggg gttgttctct tagagcttgt 1740 aactggacgg aaaccagttg accagactca gcctctagga gaagagagtt tggttgaatg 1800 ggcgcgcccg ctgcttctca aagccattga gaccggagat ttaagcgaac tgattgatac 1860 acggettgaa aagegttatg tggageatga agtetteaga atgategaga eageegetge 1920 atgtgttaga cattctggtc caaaacgccc acgcatggtt caggttgtga gagcattgga 1980 ctgcgacgga gactcgggag atattagcaa cggaatcaaa attgggcaaa gcacaactta 2040 tgactcaggg caatacaatg aagacattat gaaattcagg aaaatggcgt ttggtggtga 2100 taacagcgta gagtcaggat tgtacagtgg aaactactct gccaaaagct cttcagattt 2160 ctcagggaat gaatctgaga ctcggccttt caacaaccga cggttctgat catacaatag 2220 gtgaaagtaa c

<210> 7 <211> 2196

<212> DNA
<213> Arabidopsis thaliana
<220>
<221> CDS
<222> (1)..(2196)
<400> 7
atg tca gac tta ggc gag tce
Met Ser Asp Leu Glv Glu Se

<400> 7 atg tca gac tta ggc gag tcg ccg agt tct tca cca cca qca cca 48 Met Ser Asp Leu Gly Glu Ser Pro Ser Ser Pro Pro Ala Pro Pro get gat ace get eet eea eea gag act eea tea gaa aac tea get ett 96 Ala Asp Thr Ala Pro Pro Pro Glu Thr Pro Ser Glu Asn Ser Ala Leu cca cct gtt gat tcc tct cct cct agt cca cca gct gat tca tca 144 Pro Pro Val Asp Ser Ser Pro Pro Ser Pro Pro Ala Asp Ser Ser Ser 40 aca ccg ccg ctg tca gaa cca tcc act cct cca gat tca cag ctt 192 Thr Pro Pro Leu Ser Glu Pro Ser Thr Pro Pro Pro Asp Ser Gln Leu 50 55 cct cct tta cct tcg att ctt cct ccg cta aca gat tct cca cct cca 240 Pro Pro Leu Pro Ser Ile Leu Pro Pro Leu Thr Asp Ser Pro Pro cet tee gat tet tet eea eee gtt gat tea ace eet tet eeg eeg eea 288 Pro Ser Asp Ser Ser Pro Pro Val Asp Ser Thr Pro Ser Pro Pro Pro ccg acg tca aac gaa tct cct tct cct cca gaa gat tcc gaa aca cca 336 Pro Thr Ser Asn Glu Ser Pro Ser Pro Pro Glu Asp Ser Glu Thr Pro 100 105 cct gct cca cca aat gaa tcc aat gac aac aac cct cct ccg tct caa 384 Pro Ala Pro Pro Asn Glu Ser Asn Asp Asn Asn Pro Pro Pro Ser Gln 115 120 gat ctt caa tcg cct cct cca tcg tcg ccg tcg ccg aat gta gga ccc 432 Asp Leu Gln Ser Pro Pro Pro Ser Ser Pro Ser Pro Asn Val Gly Pro 130 135 aca aac ccg gaa tca cca ccg tta caa tct cct cca gct cca cca gca 480 Thr Asn Pro Glu Ser Pro Pro Leu Gln Ser Pro Pro Ala Pro Pro Ala 145 150 160 155 tca gat cct aca aat tca ccg cca gct tca cca tta gac cct acc aat 528 Ser Asp Pro Thr Asn Ser Pro Pro Ala Ser Pro Leu Asp Pro Thr Asn 165 cct ccc cca ata caa cca tca gga cca gcc act tct cct ccg gct aat 576

Pro Pro Pro Ile Gln Pro Ser Gly Pro Ala Thr Ser Pro Pro Ala Asn

ccc Pro	aac Asn	gct Ala 195	ccg Pro	ccg Pro	agc Ser	cca Pro	ttc Phe 200	ccc Pro	aca Thr	gta Val	cca Pro	ccc Pro 205	aaa Lys	act Thr	cct Pro	624
tct Ser	agt Ser 210	gga Gly	cct Pro	gtg Val	gtg Val	tct Ser 215	cca Pro	tct Ser	ctc Leu	aca Thr	tcc Ser 220	cct Pro	agt Ser	aaa Lys	gga Gly	672
act Thr 225	cct Pro	act Thr	cca Pro	aac Asn	caa Gln 230	ggc Gly	aat Asn	gga Gly	gat Asp	ggc Gly 235	ggt Gly	ggc Gly	ggt Gly	ggt Gly	ggc Gly 240	720
ggc Gly	tat Tyr	caa Gln	Gly 999	aag Lys 245	act Thr	atg Met	gtt Val	ggt Gly	atg Met 250	gct Ala	gta Val	gcc Ala	ggt Gly	ttc Phe 255	gca Ala	768
atc <u>I</u> le	atg Met	gcg Ala	ctt Leu 260	ata Ile	ggc Gly	gtt Val	gtg Val	ttc Phe 265	tta Leu	gtg Val	aga Arg	aga Arg	aag Lys 270	aaa Lys	aag Lys	816
	Asn				tat Tyr											864
ttc Phe	tct Ser 290	gtt Val	aaa Lys	tca Ser	gat Asp	gga Gly 295	ttc Phe	tta Leu	tac Tyr	ggt Gly	caa Gln 300	gat Asp	cca Pro	ggt Gly	aaa Lys	912
gga Gly 305	tac Tyr	tcc Ser	tct Ser	ggt Gly	cct Pro 310	aat Asn	ggt Gly	tca Ser	atg Met	tat Tyr 315	aac Asn	aat Asn	tca Ser	cag Gln	caa Gln 320	960
caa Gln	caa Gln	tcc Ser	tct Ser	atg Met 325	gga Gly	aac Asn	agt Ser	tat Tyr	ggt Gly 330	aca Thr	gct Ala	ggt Gly	ggt Gly	ggt Gly 335	tat Tyr	1008
cct Pro	cat His	cat His	caa Gln 340	atg Met	caa Gln	tca Ser	agt Ser	ggc Gly 345	aca Thr	cct Pro	gac Asp	tct Ser	gct Ala 350	ata Ile	ctc Leu	1056
					cat His											1104
caa Gln	ggc Gly 370	ttt Phe	gct Ala	cgc Arg	aaa Lys	aac Asn 375	att Ile	ctt Leu	gga Gly	gaa Glu	ggc Gly 380	gga Gly	ttt Phe	gga Gly	tgt Cys	1152
gtc Val 385	tat Tyr	aaa Lys	ggt Gly	aca Thr	ttg Leu 390	cag Gln	gat Asp	ggt Gly	aaa Lys	gtt Val 395	gtt Val	gcg Ala	gtt Val	aag Lys	cag Gln 400	1200

	aaa Lys															1248
	atc Ile															1296
	tgc Cys															1344
	caa Gln 450															1392
ato Ile 465	gct Ala	ata Ile	gga Gly	tca Ser	gcc Ala 470	aaa Lys	Gly aaa	ttg Leu	gca Ala	tat Tyr 475	ctt Leu	cac His	gaa Glu	gac Asp	tgt Cys 480	1440
	ccg Pro															1488
	gat Asp															1536
	cta Leu															1584
	ctc Leu 530															1632
	ttc Phe															1680
	aga Arg															1728
	gga Gly															1776
	gtt Val															1824
gat Asp	tta Leu	agc Ser	gaa Glu	ctg Leu	att Ile	gat Asp	aca Thr	cgg Arg	ctt Leu	gaa Glu	aag Lys	cgt Arg	tat Tyr	gtg Val	gag Glu	1872

cat His 625	gaa Glu	gtc Val	ttc Phe	aga Arg	atg Met 630	atc Ile	gag Glu	aca Thr	gcc Ala	gct Ala 635	gca Ala	tgt Cys	gtt Val	aga Arg	cat His 640	1920
tct Ser	ggt Gly	cca Pro	aaa Lys	cgc Arg 645	cca Pro	cgc Arg	atg Met	gtt Val	cag Gln 650	gtt Val	gtg Val	aga Arg	gca Ala	ttg Leu 655	gac Asp	1968
tgc Cys	gac Asp	gga Gly	gac Asp 660	tcg Ser	gga Gly	gat Asp	att Ile	agc Ser 665	aac Asn	gga Gly	atc Ile	aaa Lys	att Ile 670	Gly 999	caa Gln	2016
														aaa Lys		2064
agg Arg	aaa Lys 690	atg Met	gcg Ala	ttt Phe	ggt Gly	ggt Gly 695	gat Asp	aac Asn	agc Ser	gta Val	gag Glu 700	tca Ser	gga Gly	ttg Leu	tac Tyr	2112
agt Ser 705	gga Gly	aac Asn	tac Tyr	tct Ser	gcc Ala 710	aaa Lys	agc Ser	tct Ser	tca Ser	gat Asp 715	ttc Phe	tca Ser	G1 y 999	aat Asn	gaa Glu 720	2160
tct Ser	gag Glu	act Thr	cgg Arg	cct Pro 725	ttc Phe	aac Asn	aac Asn	cga Arg	cgg Arg 730	ttc Phe	tga					2196

<210> 8

<211> 731

<212> PRT

<213> Arabidopsis thaliana

<400> 8

Met Ser Asp Leu Gly Glu Ser Pro Ser Ser Pro Pro Ala Pro Pro 5 10 Ala Asp Thr Ala Pro Pro Pro Glu Thr Pro Ser Glu Asn Ser Ala Leu 20 25 Pro Pro Val Asp Ser Ser Pro Pro Ser Pro Pro Ala Asp Ser Ser Ser 40 45 Thr Pro Pro Leu Ser Glu Pro Ser Thr Pro Pro Pro Asp Ser Gln Leu 55 Pro Pro Leu Pro Ser Ile Leu Pro Pro Leu Thr Asp Ser Pro Pro 70 75 Pro Ser Asp Ser Ser Pro Pro Val Asp Ser Thr Pro Ser Pro Pro 85 90 Pro Thr Ser Asn Glu Ser Pro Ser Pro Pro Glu Asp Ser Glu Thr Pro 100 105 Pro Ala Pro Pro Asn Glu Ser Asn Asp Asn Pro Pro Pro Ser Gln 115

120

Asp Leu Gln Ser Pro Pro Pro Ser Ser Pro Ser Pro Asn Val Gly Pro Thr Asn Pro Glu Ser Pro Pro Leu Gln Ser Pro Pro Ala Pro Pro Ala Ser Asp Pro Thr Asn Ser Pro Pro Ala Ser Pro Leu Asp Pro Thr Asn Pro Pro Pro Ile Gln Pro Ser Gly Pro Ala Thr Ser Pro Pro Ala Asn Pro Asn Ala Pro Pro Ser Pro Phe Pro Thr Val Pro Pro Lys Thr Pro Ser Ser Gly Pro Val Val Ser Pro Ser Leu Thr Ser Pro Ser Lys Gly Thr Pro Thr Pro Asn Gln Gly Asn Gly Asp Gly Gly Gly Gly Gly Gly Tyr Gln Gly Lys Thr Met Val Gly Met Ala Val Ala Gly Phe Ala Ile Met Ala Leu Ile Gly Val Val Phe Leu Val Arg Arg Lys Lys Arg Asn Ile Asp Ser Tyr Asn His Ser Gln Tyr Leu Pro His Pro Asn Phe Ser Val Lys Ser Asp Gly Phe Leu Tyr Gly Gln Asp Pro Gly Lys Gly Tyr Ser Ser Gly Pro Asn Gly Ser Met Tyr Asn Asn Ser Gln Gln Gln Gln Ser Ser Met Gly Asn Ser Tyr Gly Thr Ala Gly Gly Gly Tyr Pro His His Gln Met Gln Ser Ser Gly Thr Pro Asp Ser Ala Ile Leu Gly Ser Gly Gln Thr His Phe Ser Tyr Glu Glu Leu Ala Glu Ile Thr Gln Gly Phe Ala Arg Lys Asn Ile Leu Gly Glu Gly Gly Phe Gly Cys Val Tyr Lys Gly Thr Leu Gln Asp Gly Lys Val Val Ala Val Lys Gln Leu Lys Ala Gly Ser Gly Gln Gly Asp Arg Glu Phe Lys Ala Glu Val Glu Ile Ile Ser Arg Val His His Arg His Leu Val Ser Leu Val Gly Tyr Cys Ile Ser Asp Gln His Arg Leu Leu Ile Tyr Glu Tyr Val Ser Asn Gln Thr Leu Glu His His Leu His Glu Trp Ser Lys Arg Val Arg Ile Ala Ile Gly Ser Ala Lys Gly Leu Ala Tyr Leu His Glu Asp Cys His Pro Lys Ile Ile His Arg Asp Ile Lys Ser Ala Asn Ile Leu Leu Asp Asp Glu Tyr Glu Ala Gln Ala Ile Met Lys Ser Ser Phe Ser Leu Asn Leu Ser Tyr Asp Cys Lys Val Leu Val Ala Asp Phe Gly Leu Ala Arg Leu Asn Asp Thr Thr Gln Thr His Val Ser Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ser Ser Gly Lys Leu Thr 

```
Asp Arg Ser Asp Val Phe Ser Phe Gly Val Val Leu Leu Glu Leu Val
                565
                                     570
Thr Gly Arg Lys Pro Val Asp Gln Thr Gln Pro Leu Gly Glu Glu Ser
            580
                                 585
                                                     590
Leu Val Glu Trp Ala Arg Pro Leu Leu Leu Lys Ala Ile Glu Thr Gly
                             600
Asp Leu Ser Glu Leu Ile Asp Thr Arg Leu Glu Lys Arg Tyr Val Glu
                        615
                                             620
His Glu Val Phe Arg Met Ile Glu Thr Ala Ala Ala Cys Val Arg His
                    630
                                         635
Ser Gly Pro Lys Arg Pro Arg Met Val Gln Val Val Arg Ala Leu Asp
                645
                                     650
Cys Asp Gly Asp Ser Gly Asp Ile Ser Asn Gly Ile Lys Ile Gly Gln
            660
                                . 665
Ser Thr Thr Tyr Asp Ser Gly Gln Tyr Asn Glu Asp Ile Met Lys Phe
                            680
Arg Lys Met Ala Phe Gly Gly Asp Asn Ser Val Glu Ser Gly Leu Tyr
                        695
                                             700
Ser Gly Asn Tyr Ser Ala Lys Ser Ser Ser Asp Phe Ser Gly Asn Glu
                    710
                                         715
Ser Glu Thr Arg Pro Phe Asn Asn Arg Arg Phe
                725
```

```
<210> 9
<211> 1939
<212> DNA
<213> Arabidopsis thaliana
```

```
₹400> 9
gaaaattttg atctccgatg gcttcttctc ctgaatctgc tcctccaaca aactccacct 60
cttctccatc tccaccgtct aataccaatt caaccacctc ttctccgccg gctccgtctc 120
ctccttctcc tacacctcct caaggagact catcatcatc gccacctcct gattccacat 180
ctccaccage tecacaaget ectaaceete etaatteete taataaetet eetteeete 240
cgtcacaggg cggtggagga gaaagaggaa atggaggaaa caatggtggc aatgatactc 300
caccgtcacg cggctctcct ccttctcctc cttctaggag taatggagat aatggtggta 360
gcagatcatc gccaccagga gacactggag gctctcgctc agacaaccct ccttctagcg 420
gaggaagcag tggaggaggt ggaggtggaa gaagtaatac gaatacagcg atcatagttg 480
gtgtattagt cggagctgga cttttgatga tcgttcttat tattgtgtgt cttagacgca 540
aaaagaagag aaaagactcc ttctaccctg aacccatgaa aggaaaccaa tatcaatact 600
atggaaacaa caacaacaac aatgcttcac agaattatcc gaattggcac ctaaattcac 660
aaggccaaaa ccaacaatct actggtggtt ggggaggcgg tggaccatca ccgcctcctc 720
ctccgcggat gcctacaagc ggagaagatt cttccatgta ctcaggccca tcacgcccag 780
ttttacctcc tccttcgcct gctctagccc tcggattcaa caagagcact tttacttacc 840
aagagettge ggetgeaaca ggagggttta eggatqetaa eettttqqqa eaqqqaqqat 900
ttgggtatgt ccataaagga gtcttgccta gcgggaaaga agtagcagtt aagagtttaa 960
agcgggtag cggacaagga gagagggagt ttcaagctga ggtcgatatc attagccgtg 1020°
tgeateateg gtatettgtt tetttggttg gatattgeat agetgatgga eagaggatgt 1080
tggtttatga gtttgttcct aacaaaactt tggaatatca tcttcatggg aaaaatcttc 1140
cggtaatgga gttctccact aggttgcgta tcgccttagg tgctgcgaaa ggactcgctt 1200
accttcacga agactgccat cctcggatca ttcaccgcga catcaagtct gcaaatattc 1260
tcttggactt caactttgat gctatggtgg ctgattttgg attagctaag ttaacatctq 1320
ataacaacac tcatgtatct actcgtgtga tgggaacttt cggatatcta gctccaqaat 1380
```

```
tggaacttat aactggaaaa cgaccggttg ataatagcat caccatggac gacaccttag 1500
tagattgggc tcggcctctt atggctcgcg cgctagaaga tggaaacttt aatgagctcg 1560
Eagatgcgag gcttgaaggc aactacaacc cgcaagaaat ggctcgaatg gtgacttgtg 1620
cgctgctag cattcgtcat tcggggcgta aacgtccaaa gatgagccag atagtaagag 1680
cgttagaagg agaagtgtcc ttagatgctt taaacgaagg tgtgaagcca ggacacagta 1740
acgtttacgg gtcattggga gcaagctcgg attatagtca gacatcttac aatgcagaca 1800
tgaagaaatt cagacagata gctttgtcga gccaagaatt cccagtcagt gactgtgaag 1860
gaacatctag taatgattct agagatatgg gaactaagag ccctactcct ccaaaatgag 1920
atcqaatcaa tqattctqt
<210> 10
<211> 1902
<212> DNA
<213> Arabidopsis thaliana
<220>
<221> CDS
<222> (1)..(1902)
<sup>2</sup>400> 10
atg gct tct tct cct gaa tct gct cct cca aca aac tcc acc tct tct
Met Ala Ser Ser Pro Glu Ser Ala Pro Pro Thr Asn Ser Thr Ser Ser
  1
                                    10
                                                       15
cca tct cca ccg tct aat acc aat tca acc acc tct tct ccg ccg gct
                                                                96
Pro Ser Pro Pro Ser Asn Thr Asn Ser Thr Thr Ser Ser Pro Pro Ala
ccg tct cct tct cct aca cct cct caa gga gac tca tca tcg
                                                                144
Pro Ser Pro Pro Ser Pro Thr Pro Pro Gln Gly Asp Ser Ser Ser
                            40
cca cct cct gat tcc aca tct cca cca gct cca caa gct cct aac cct
                                                                192
Pro Pro Pro Asp Ser Thr Ser Pro Pro Ala Pro Gln Ala Pro Asn Pro
     50
                        55
ect aat tee tet aat aac tet eet tee eet eeg tea eag gge ggt gga
                                                                240
Pro Asn Ser Ser Asn Asn Ser Pro Ser Pro Pro Ser Gln Gly Gly
 65
                    70
gga gaa aga gga aat gga gga aac aat ggt ggc aat gat act cca ccg
                                                                288
Gly Glu Arg Gly Asn Gly Gly Asn Gly Gly Asn Asp Thr Pro Pro
tca cgc ggc tct cct cct tct cct tct agg agt aat gga gat aat
                                                                336
Ser Arg Gly Ser Pro Pro Ser Pro Pro Ser Arg Ser Asn Gly Asp Asn
           100
                               105
                                                  110
ggt ggt agc aga tca tcg cca cca gga gac act gga ggc tct cgc tca
                                                                384
Gly Gly Ser Arg Ser Ser Pro Pro Gly Asp Thr Gly Gly Ser Arg Ser
```

aac caa caa tct act ggt ggt 672 Asn Gln Gln Ser Thr Gly Gly 220 cct cct ccg cgg atg cct aca 720 Pro Pro Pro Arg Met Pro Thr	235 240 gc cca tca cgc cca gtt tta 768 ly Pro Ser Arg Pro Val Leu
	sn Gln Gln Ser Thr Gly Gly 220 et eet eeg egg atg eet aca 72
235 240 gc cca tca cgc cca gtt tta 768 Ly Pro Ser Arg Pro Val Leu	
235  240  c cca tca cgc cca gtt tta 768 y Pro Ser Arg Pro Val Leu 0 255  a ttc aac aag agc act ttt 816 y Phe Asn Lys Ser Thr Phe 270	y Phe Asn Lys Ser Thr Phe 270
cca tca cgc cca gtt tta 768 Pro Ser Arg Pro Val Leu 255  ttc aac aag agc act ttt 816 Phe Asn Lys Ser Thr Phe 270  ggg ttt acg gat gct aac 864 Gly Phe Thr Asp Ala Asn 285	Phe Asn Lys Ser Thr Phe 270  ggg ttt acg gat gct aac 864 Gly Phe Thr Asp Ala Asn 285
tca cgc cca gtt tta 768 Ser Arg Pro Val Leu 255  aac aag agc act ttt 816 Asn Lys Ser Thr Phe 270  ttt acg gat gct aac 864 Phe Thr Asp Ala Asn 285  aaa gga gtc ttg cct 912 Lys Gly Val Leu Pro 300	Asn Lys Ser Thr Phe 270  ttt acg gat gct aac 864 Phe Thr Asp Ala Asn 285  aaa gga gtc ttg cct 912 Lys Gly Val Leu Pro 300
240  cca tca cgc cca gtt tta 768  cro Ser Arg Pro Val Leu 255  ctc aac aag agc act ttt 816  che Asn Lys Ser Thr Phe 270  ggg ttt acg gat gct aac 864  Gly Phe Thr Asp Ala Asn 285  cat aaa gga gtc ttg cct 912  dis Lys Gly Val Leu Pro 300  aaa gcg ggt agc gga caa 960  cys Ala Gly Ser Gly Gln	Phe Asn Lys Ser Thr Phe 270  ggg ttt acg gat gct aac 864 Gly Phe Thr Asp Ala Asn 285  cat aaa gga gtc ttg cct 912 His Lys Gly Val Leu Pro 300  aaa gcg ggt agc gga caa 960 Lys Ala Gly Ser Gly Gln
ggc cca tca cgc cca gtt tta 768 Gly Pro Ser Arg Pro Val Leu 255  gga ttc aac aag agc act ttt 816 Gly Phe Asn Lys Ser Thr Phe 270  gga ggg ttt acg gat gct aac 864 Gly Gly Phe Thr Asp Ala Asn 285  gtc cat aaa gga gtc ttg cct 912 Al His Lys Gly Val Leu Pro 300  cta aaa gcg ggt agc gga caa 960	Iy Phe Asn Lys Ser Thr Phe 270  ga ggg ttt acg gat gct aac 864  ly Gly Phe Thr Asp Ala Asn 285  cc cat aaa gga gtc ttg cct 912  al His Lys Gly Val Leu Pro 300  ca aaa gcg ggt agc gga caa 960  eu Lys Ala Gly Ser Gly Gln 315  at atc att agc cgt gtg cat 1006  sp Ile Ile Ser Arg Val His

•

•												,				
His	Arg	Tyr	Leu 340	Val	Ser	Leu	Val	Gly 345	Tyr	Cys	Ile	Ala	Asp 350	Gly	Gln	
	_	_	-				_					_	_	tat Tyr		1104
														ttg Leu		1152
	_			-					_				_	gac Asp	_	1200
						_	_		_		_			ctc Leu 415	_	1248
														aag Lys		1296
														act Thr		1344
														gag Glu		1392
														act Thr		1440
														gta Val 495		1488
														ttt Phe		1536
														gaa Glu		1584
														Gly ggg		1632
														gaa Glu		1680

•

tcc Ser	tta Leu	gat Asp	gct Ala	tta Leu 565	aac Asn	gaa Glu	ggt Gly	gtg Val	aag Lys 570	cca Pro	gga Gly	cac His	agt Ser	aac Asn 575	gtt Val	1728
tac Tyr	Gly aaa	tca Ser	ttg Leu 580	gga Gly	gca Ala	agc Ser	tcg Ser	gat Asp 585	tat Tyr	agt Ser	cag Gln	aca Thr	tct Ser 590	tac Tyr	aat Asn	1776
gca Ala	gac Asp	atg Met 595	aag Lys	aaa Lys	ttc Phe	aga Arg	cag Gln 600	ata Ile	gct Ala	ttg Leu	tcg Ser	agc Ser 605	caa Gln	gaa Glu	ttc Phe	1824
cca Pro	gtc Val 610	agt Ser	gac Asp	tgt Cys	gaa Glu	gga Gly 615	aca Thr	tct Ser	agt Ser	aat Asn	gat Asp 620	tct Ser	aga Arg	gat Asp	atg Met	1872
					act Thr 630				tga							1902

<210> 11

<211> 633

<212> PRT <213> Arabidopsis thaliana

<400> 11

Met 1	Ala	Ser	Ser	Pro 5	Glu	Ser	Ala	Pro	Pro 10	Thr	Asn	Ser	Thr	Ser 15	Ser
Pro	Ser	Pro	Pro 20	Ser	Asn	Thr	Asn	Ser 25	Thr	Thr	Ser	Ser	Pro 30	Pro	Ala
Pro	Ser	Pro 35	Pro	Ser	Pro	Thr	Pro 40	Pro	Gln	Gly	Asp	Ser 45	Ser	Ser	Ser
Pro	Pro 50	Pro	Asp	Ser	Thr	Ser 55	Pro	Pro	Ala	Pro	Gln 60	Ala	Pro	Asn	Pro
Pro 65	Asn	Ser	Ser	Asn	Asn 70	Ser	Pro	Ser	Pro	Pro 75	Ser	Gln	Gly	Gly	Gly 80
Gly	Glu	Arg	Gly	Asn 85	Gly	Gly	Asn	Asn	Gly 90	Gly	Asn	Asp	Thr	Pro 95	Pro
Ser	Arg	Gly	Ser 100	Pro	Pro	Ser	Pro	Pro 105	Ser	Arg	Ser	Asn	Gly 110	Asp	Asn
Gly	Gly	Ser 115	Arg	Ser	Ser	Pro	Pro 120	Gly	Asp	Thr	Gly	Gly 125	Ser	Arg	Ser
Asp	Asn 130	Pro	Pro	Ser	Ser	Gly 135	Gly	Ser	Ser	Gly	Gly 140	Gly	Gly	Gly	Gly
Arg 145	Ser	Asn	Thr	Asn	Thr 150	Ala	Ile	Ile	Val	Gly 155	Val	Leu	Val	Gly	Ala 160
Gly	Leu	Leu	Met	Ile 165	Val	Leu	Ile	Ile	Val 170	Cys	Leu	Arg	Arg	Lys 175	Lys
Lys	Arg	Lys	Asp 180	Ser	Phe	Tyr	Pro	Glu 185	Pro	Met	Lys	Gly	Asn 190	Gln	Tyr
Gln	Tyr	Tyr 195	Gly	Asn	Asn	Asn	Asn 200	Asn	Asn	Ala	Ser	Gln 205	Asn	Tyr	Pro

Asn Trp His Leu Asn Ser Gln Gly Gln Asn Gln Gln Ser Thr Gly Gly Trp Gly Gly Gly Pro Ser Pro Pro Pro Pro Pro Arg Met Pro Thr Ser Gly Glu Asp Ser Ser Met Tyr Ser Gly Pro Ser Arg Pro Val Leu Pro Pro Pro Ser Pro Ala Leu Ala Leu Gly Phe Asn Lys Ser Thr Phe Thr Tyr Gln Glu Leu Ala Ala Ala Thr Gly Gly Phe Thr Asp Ala Asn Leu Leu Gly Gln Gly Gly Phe Gly Tyr Val His Lys Gly Val Leu Pro Ser Gly Lys Glu Val Ala Val Lys Ser Leu Lys Ala Gly Ser Gly Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Asp Ile Ile Ser Arg Val His His Arg Tyr Leu Val Ser Leu Val Gly Tyr Cys Ile Ala Asp Gly Gln Arg Met Leu Val Tyr Glu Phe Val Pro Asn Lys Thr Leu Glu Tyr His Leu His Gly Lys Asn Leu Pro Val Met Glu Phe Ser Thr Arg Leu Arg Ile Ala Leu Gly Ala Ala Lys Gly Leu Ala Tyr Leu His Glu Asp Cys His Pro Arg Ile Ile His Arg Asp Ile Lys Ser Ala Asn Ile Leu Leu Asp Phe Asn Phe Asp Ala Met Val Ala Asp Phe Gly Leu Ala Lys Leu Thr Ser Asp Asn Asn Thr His Val Ser Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ser Ser Gly Lys Leu Thr Glu Lys Ser Asp Val Phe Ser Tyr Gly Val Met Leu Leu Glu Leu Ile Thr Gly Lys Arg Pro Val Asp Asn Ser Ile Thr Met Asp Asp Thr Leu Val Asp Trp Ala Arg Pro Leu Met Ala Arg Ala Leu Glu Asp Gly Asn Phe Asn Glu Leu Ala Asp Ala Arg Leu Glu Gly Asn Tyr Asn Pro Gln Glu Met Ala Arg Met Val Thr Cys Ala Ala Ala Ser Ile Arg His Ser Gly Arg Lys Arg Pro Lys Met Ser Gln Ile Val Arg Ala Leu Glu Gly Glu Val Ser Leu Asp Ala Leu Asn Glu Gly Val Lys Pro Gly His Ser Asn Val Tyr Gly Ser Leu Gly Ala Ser Ser Asp Tyr Ser Gln Thr Ser Tyr Asn Ala Asp Met Lys Lys Phe Arg Gln Ile Ala Leu Ser Ser Gln Glu Phe Pro Val Ser Asp Cys Glu Gly Thr Ser Ser Asn Asp Ser Arg Asp Met Gly Thr Lys Ser Pro Thr Pro Pro Lys 

```
<210> 12
<211> 2104
<212> DNA
<213> Arabidopsis thaliana
<400> 12
tccaccgttt gagaaaccct aataacaaca ttcaaaatgg cggactcacc ggtggattca 60
tetectgece etgaaacete aaatgggaca ecacegteaa aeggaacate geegtetaat 120
gagtcatcgc cgccaacacc accttcttca ccaccaccat catcaatatc tqctcctccq 180
ccagatatct ccgcttcttt ttcaccgccg cctgcaccac caacgcaaga aacgtcacct 240
cctacatete egtecteate geegeetgtt gtagetaate egteacegea gaeteeagag 300
aatcettete cacetgeace tgaaggetea acteetgtaa egecacetge accaceacaa 360
acaccgtcga accaatcacc ggaaagacca actcctcctt ctcctggtgc caatgatgac 420
cgaaacagaa ccaatggcgg aaacaacaac agagacggct ccacaccatc accaccqtcq 480
tcagggaaca gaacttccgg tgacggtggc tcaccttcac cacctcggtc gataagccct 540
cctcagaata gtggagattc agactcatca tcggggcttt tgcttctact tqcaqtqtqt 600
atttgcatct gttgcaacag gaagaagaag aagaaatctc ctcaqqtcaa ccacatqcac 660
factacaata acaateetta tggaggagea eeeteaggta atggtggtta ttacaaggga 720
Acaceteaag ateatgtggt gaatatgget ggteaaggag gtgggaattg gggteeaeag 780
caacctgtgt ctggtcctca cagtgatgct tccaacttaa ccggtcgaac tgctataccg 840
tcacctcaag ctgcaactct tggtcacaac caaaqcactt tcacatacqa tgaactgtcc 900
attgcaacag aaggtttege teagteaaat ttgetaggae aaggaggatt tgggtatgtt 960
cataaaggag ttctgcctag tggcaaagaa gttgcagtga agagtcttaa acttggaagt 1020
ggacaagggg aacgcgagtt tcaagcagag gttgatatca ttagccgtgt ccatcatcgt 1080
catctcgttt ctcttgttgg atattgcatc tctggtggtc aaagactttt ggtttatgag 1140
tttataccta acaacactct tgaatttcat cttcatggaa agggtcgtcc ggttttggat 1200
tggcctacaa gagtgaagat tgcattggga tcagctagag gccttgcata tttgcatqaa 1260
gactgtaaga aaatctttat ctcacatatt tgcatcagtc accctcgcat tatccacaqa 1320
gatatcaaag ctgcaaacat tcttcttgat ttcagttttg agaccaaggt ggcagatttt 1380
ggattggcta agctatctca agacaactat actcatgtct ccactcgcgt catgggaact 1440
tttggatact tagctccaga gtatgcatca agcggaaagt tatccgacaa atctgatgtt 1500
tteteatttg gagtaatget tettgagete ataaceggaa gaeeteetet ggatetaaet 1560
ggagaaatgg aagatagctt ggtagattgg gcaaggcctt tgtgtttgaa agcagctcaa 1620
gatggagatt acaaccaatt ggctgatcca cgtctagagc taaactacag tcatcaaqag 1680
åtggttcaaa tggcttcttg tgcagctgca gcaatcagac attcagcaag aagacggcct 1740
agatgagcc aggttcaaaa actcatacca cttgttggtt ctattattgt acgagcacta 1800
gaaggagata tgtcaatgga tgatctaagt qaqqqaacaa qaccaqqaca aaqcacqtac 1860
ttgagccccg ggagcgtgag ctcagagtat gacgcaagct cgtacacggc agacatgaaa 1920
aagttcaaga aactggcgtt agagaataaa gaatatcaaa gcagtgaata tgqtggaaca 1980
agtgagtatg gcttaaaccc ttctgcttca agtagtgaag aaatgaatag aggctcaatg 2040
aaacgcaatc ctcagctttg aaagaagaga caacacttgt cataatattt cagttttctt 2100
ctct
                                                                   2104
<210> 13
<211> 2025
<212> DNA
<213> Arabidopsis thaliana
<220>
₹221> CDS
```

<222> (1)..(2025)

atg	Ala	gac	: tca Ser	. ccg Pro	Val	gat Asp	tca Ser	tct Ser	cct Pro	Ala	c cct Pro	gaa Glu	acc Thr	tca Ser 15	aat Asn	48
Gly	aca Thr	cca Pro	ccg Pro 20	Ser	aac Asn	gga Gly	aca Thr	tcg Ser 25	Pro	tct Ser	aat Asn	gag Glu	tca Ser 30	Ser	ccg Pro	96
cca Pro	aca Thr	cca Pro 35	Pro	tct Ser	tca Ser	cca Pro	cca Pro 40	cca Pro	tca Ser	tca Ser	ata Ile	tct Ser 45	gct Ala	cct Pro	ccg Pro	144
cca Pro	gat Asp 50	Ile	tcc Ser	gct Ala	tct Ser	ttt Phe 55	tca Ser	ccg Pro	ccg Pro	cct Pro	gca Ala 60	cca Pro	cca Pro	acg Thr	caa Gln	192
gaa Glu 65	acg Thr	tca Ser	cct Pro	cct Pro	aca Thr 70	tct Ser	ccg Pro	tcc Ser	tca Ser	tcg Ser 75	Pro	cct Pro	gtt Val	gta Val	gct Ala 80	240
aat Asn	ccg Pro	tca Ser	ccg Pro	cag Gln 85	act Thr	cca Pro	gag Glu	aat Asn	cct Pro 90	tct Ser	cca Pro	cct Pro	gca Ala	cct Pro 95	gaa Glu	288
ggc Gly	tca Ser	act Thr	cct Pro 100	gta Val	acg Thr	cca Pro	cct Pro	gca Ala 105	cca Pro	cca Pro	caa Gln	aca Thr	ccg Pro 110	tcg Ser	aac Asn	336
caa Gln	tca Ser	ccg Pro 115	gaa Glu	aga Arg	cca Pro	act Thr	cct Pro 120	cct Pro	tct Ser	cct Pro	ggt Gly	gcc Ala 125	aat Asn	gat Asp	gac Asp	384
cga Arg ,	aac Asn 130	aga Arg	acc Thr	aat Asn	ggc Gly	gga Gly 135	aac Asn	aac Asn	aac Asn	aga Arg	gac Asp 140	ggc Gly	tcc Ser	aca Thr	cca Pro	432
tca Ser 145	cca Pro	ccg Pro	tcg Ser	tca Ser	999 Gly 150	aac Asn	aga Arg	act Thr	tcc Ser	ggt Gly 155	gac Asp	ggt Gly	ggc Gly	tca Ser	cct Pro 160	480
sca Ser	cca Pro	cct Pro	cgg Arg	tcg Ser 165	ata Ile	agc Ser	cct Pro	cct Pro	cag Gln 170	aat Asn	agt Ser	gga Gly	gat Asp	tca Ser 175	gac Asp	528
tca Ser	tca Ser	tcg Ser	999 Gly 180	ctt Leu	ttg Leu	ctt Leu	cta Leu	ctt Leu 185	gca Ala	gtg Val	tgt Cys	att Ile	tgc Cys 190	atc Ile		576
tgc Cys	Asn	agg Arg 195	aag Lys	aag Lys	aag Lys	aag Lys	aaa Lys 200	tct Ser	cct Pro	cag Gln	Val	aac Asn 205	cac His	atg Met	cac His	624

	tac Tyr 210															672
	tac Tyr															720
	ggt Gly															768
	gct Ala															816
	act Thr															864
	gca Ala 290															912
	gjå aaa															960
	aag Lys															1008
	gag Glu															1056
	gtt Val															1104
	ata Ile 370															1152
	gtt Val															1200
	ggc Gly															1248
cat	att	tgc	atc	agt	cac	cct	cgc	att	atc	cac	aga	gat	atc	aaa	gct	1296

His	Ile	Cys	Ile 420	Ser	His	Pro	Arg	Ile 425	Ile	His	Arg	Asp	Ile 430	Lys	Ala	
					gat Asp											1344
äga					tct Ser											1392
					gga Gly 470											1440
					tct Ser											1488
					aga Arg											1536
					tgg Trp											1584
					caa Gln											1632
					gtt Val 550											1680
_		_			aga Arg	_	_							_		1728
ata 11e	cca Pro	ctt Leu	gtt Val 580	ggt Gly	tct Ser	att Ile	att Ile	gta Val 585	cga Arg	gca Ala	cta Leu	gaa Glu	gga Gly 590	gat Asp	atg Met	1776
					agt Ser											1824
_	_			_	gtg Val	_				_	_	_	_		_	1872
					ttc Phe 630											1920

caa Gln	_	_	_			_	gag Glu 650				1968
							ggc Gly				2016
cag Gln	ctt Leu	tga									2025

<210> 14

<211> 674

<212> PRT

<213> Arabidopsis thaliana

<400> 14

Met 1	Ala	Asp	Ser	Pro 5	Val	Asp	Ser	Ser	Pro 10	Ala	Pro	Glu	Thr	Ser 15	Asn
Gly	Thr	Pro	Pro 20	Ser	Asn	Gly	Thr	Ser 25	Pro	Ser	Asn	Glu	Ser 30	Ser	Pro
Pro	Thr	Pro 35	Pro	Ser	Ser	Pro	Pro 40	Pro	Ser	Ser	Ile	Ser 45	Ala	Pro	Pro
Pro	Asp 50	Ile	Ser	Ala	Ser	Phe 55	Ser	Pro	Pro	Pro	Ala 60	Pro	Pro	Thr	Gln
Glu 65	Thr	Ser	Pro	Pro	Thr 70	Ser	Pro	Ser	Ser	Ser 75	Pro	Pro	Val	Val	Ala 80
Asn	Pro	Ser	Pro	Gln 85	Thr	Pro	Glu	Asn	Pro 90	Ser	Pro	Pro	Ala	Pro 95	Glu
Gly	Ser	Thr	Pro 100	Val	Thr	Pro	Pro	Ala 105	Pro	Pro	Gln	Thr	Pro 110	Ser	Asn
Gln	Ser	Pro 115	Glu	Arg	Pro	Thr	Pro 120	Pro	Ser	Pro	Gly	Ala 125	Asn	Asp	Asp
Arg	Asn 130	Arg	Thr	Asn	Gly	Gly 135	Asn	Asn	Asn	Arg	Asp 140	Gly	Ser	Thr	Pro
Ser 145	Pro	Pro	Ser	Ser	Gly 150	Asn	Arg	Thr	Ser	Gly 155	Asp	Gly	Gly	Ser	Pro 160
Ser	Pro	Pro	Arg	Ser 165	Ile	Ser	Pro	Pro	Gln 170	Asn	Ser	Gly	Asp	Ser 175	Asp
Ser	Ser	Ser	Gly 180	Leu	Leu	Leu	Leu	Leu 185	Ala	Val	Cys	Ile	Cys 190	Ile	Cys
Cys	Asn	Arg 195	Lys	Lys	Lys	Lys	Lys 200	Ser	Pro	Gln	Val	Asn 205	His	Met	His
Tyr	Tyr 210	Asn	Asn	Asn	Pro	Tyr 215	Gly	Gly	Ala	Pro	Ser 220	Gly	Asn	Gly	Gly
Tyr 225	Tyr	Lys	Gly	Thr	Pro 230	Gln	Asp	His	Val	Val 235	Asn	Met	Ala	Gly	Gln 240
Gly	Gly	Gly	Asn	Trp 245	Gly	Pro	Gln	Gln	Pro 250	Val	Ser	Gly	Pro	His 255	Ser
Asp	Ala	Ser	Asn 260	Leu	Thr	Gly	Arg	Thr 265	Ala	Ile	Pro	Ser	Pro 270	Gln	Ala

Ala Thr Leu Gly His Asn Gln Ser Thr Phe Thr Tyr Asp Glu Leu Ser Ile Ala Thr Glu Gly Phe Ala Gln Ser Asn Leu Leu Gly Gln Gly Gly he Gly Tyr Val His Lys Gly Val Leu Pro Ser Gly Lys Glu Val Ala Val Lys Ser Leu Lys Leu Gly Ser Gly Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Asp Ile Ile Ser Arg Val His His Arg His Leu Val Ser Leu Val Gly Tyr Cys Ile Ser Gly Gly Gln Arg Leu Leu Val Tyr Glu Phe Ile Pro Asn Asn Thr Leu Glu Phe His Leu His Gly Lys Gly Arg Pro Val Leu Asp Trp Pro Thr Arg Val Lys Ile Ala Leu Gly Ser Ala Arg Gly Leu Ala Tyr Leu His Glu Asp Cys Lys Lys Ile Phe Ile Ser His Ile Cys Ile Ser His Pro Arg Ile Ile His Arg Asp Ile Lys Ala Ala Asn Ile Leu Leu Asp Phe Ser Phe Glu Thr Lys Val Ala Asp Phe Gly Leu Ala Lys Leu Ser Gln Asp Asn Tyr Thr His Val Ser Thr Arg Val Met Gly Thr Phe Gly Tyr Leu Ala Pro Glu Tyr Ala Ser Ser Gly Lys Leu Ser Asp Lys Ser Asp Val Phe Ser Phe Gly Val Met Leu Leu Glu Leu Ile Thr Gly Arg Pro Pro Leu Asp Leu Thr Gly Glu Met Glu Asp Ser Leu Val Asp Trp Ala Arg Pro Leu Cys Leu Lys Ala Ala Gln Asp Gly Asp Tyr Asn Gln Leu Ala Asp Pro Arg Leu Glu Leu Asn Tyr Ser His Gln Glu Met Val Gln Met Ala Ser Cys Ala Ala Ala Ile erg His Ser Ala Arg Arg Pro Lys Met Ser Gln Val Gln Lys Leu Île Pro Leu Val Gly Ser Ile Ile Val Arg Ala Leu Glu Gly Asp Met Ser Met Asp Asp Leu Ser Glu Gly Thr Arg Pro Gly Gln Ser Thr Tyr Leu Ser Pro Gly Ser Val Ser Ser Glu Tyr Asp Ala Ser Ser Tyr Thr Ala Asp Met Lys Lys Phe Lys Lys Leu Ala Leu Glu Asn Lys Glu Tyr Gln Ser Ser Glu Tyr Gly Gly Thr Ser Glu Tyr Gly Leu Asn Pro Ser Ala Ser Ser Glu Glu Met Asn Arg Gly Ser Met Lys Arg Asn Pro Gln Leu

```
<210> 15
<211> 2261
<212> DNA
<213> Arabidopsis thaliana
<400> 15
catacatcac aaaacggcat cgttttgatg tcgctctctc cttcttcctc gccggctccq 60
gcaacttccc cgcctgccat gtcattacct ccggcagatt ccgtacctga cacgtcatca 120
cctccagctc ctcctttgtc tcctcttccc ccaccattga gctctcctcc gccgttgcct 180
tcaccaccgc ctctctccgc tcccaccgct tccccaccgc ctcttccggt tgaatcccca 240
egteteete etatagaate accaeegeet eetetaetgg aateaeetee teeteeteeg 300
ttggaatete catetecace gtetecteae gteteagete etteeggtte accgecatta 360
ccetteette cegecaaace tteteegeeg cettetteac eteceteega gacagtteeg 420
ccgggaaata cgatttctcc accacctcgt tcacttccct ccgaatcaac cccqccqqtq 480
aacacagett etectecace gecateteet eetegeegee gtagtggeee taageetteg 540
tttcctcctc ccatcaattc ttctccacca aatccttctc cgaacactcc gtcactccca 600
gaaacttete etecacetaa accacegete teaaegaege cattteeete eteateeaet 660
cccccgccta agaagtcccc tgcagcagta actcttcctt tctttgggcc agcgggccaa 720
ttaccggatg ggaccgtagc acctcctatt gggcctgtta tagagcccaa gacgagtcca 780
gccgaatcaa tateteeggg aacgeeacag eeactggtte egaagagtet acetgtaacg 840
acgtcgtatc accgatcatc cgccggattc ttatttggcg gtgtaatcgt tggagctctt 900
ctactaattc tgttaggtct tctctttgtc ttctacagag ctaccagaaa tagaaataac 960
aacagcagct ctgctcatca tcaatccaaa actccctcaa aagttcaaca tcatcggggc 1020
ggtaatgctg gtacgaacca ggcacatgtt atcacaatgc caccaccaat ccatqctaaa 1080
tatatatcta gtggaggttg tgatacgaag gagaacaatt ctgttgcgaa aaacatttca 1140
atgccatctg gaatgttctc ctacgaagaa ctttcaaaag caactggtgg attttcagag 1200
gagaaccttt tgggagaagg cggtttcgga tatgttcaca aaggagtgtt gaaaaacqqq 1260
acagaagttg cggtgaagca gctgaagatt gggagctatc aaggggaaag agaattccaa 1320
gctgaggttg acacaatcag tagggttcat cataagcacc tcgtttcatt ggttggttat 1380
tgcgttaatg gagataaaag actcttggtt tacgagtttg ttcctaaaga taccttggag 1440
ttccacttgc atgagaacag aggaagcgtg ttggaatggg aaatgaggct caggattgct 1500
gtaggagcag caaaaggatt agcttatctt catgaggatt gcagtccaac tataattcac 1560
cgtgatatca aagcagctaa tatccttcta gattccaaat ttgaggcaaa ggtctctgac 1620
tttggactag ccaagttttt ctcagacacc aattcatcat tcactcatat ctctactcga 1680
gtggtaggaa ctttcggata catggctcca gaatacgcgt ccagtggtaa agtaactgat 1740
aaatcagatg tatattcctt tggggtcgtg cttctagaac tcatcactgg acgtccatca 1800
attttcgcca aagattcttc cacaaaccag agtttagtag actgggcgag gccattgctt 1860
acgaaagcaa tctctggaga aagttttgac tttcttgtag actcaaggtt ggagaagaat 1920
tacgatacaa ctcagatggc aaacatggct gcttgtgctg ctgcttgcat acgccaatca 1980
gcttggcttc ggcctagaat gagccaggta gtacgtgctc ttgaaggcga ggtggccctg 2040
agaaaggtcg aagagactgg gaatagcgtg acctatagct cttctgaaaa cccgaatgac 2100
atcacaccac ggtatggaac aaataagagg agattcgaca caggttcaag cgatggttac 2160
acttcagaat atggagttaa cccttctcag tcgagcagtg aacatcaaca ggtgaatact 2220
agttcacag gttcaatagg gcaagtttca ccacaattat t
                                                                  2261
<210> 16
<211> 2196
<212> DNA
<213> Arabidopsis thaliana
<220>
<221> CDS
<222> (1)..(2196)
```

tr Fr																
2	0> 1	6														
atg	tcg	ctc	tct Ser	cct Pro 5	Ser	tcc Ser	tcg Ser	ccg Pro	gct Ala 10	ccg Pro	gca Ala	act Thr	tcc Ser	ccg Pro 15	cct Pro	48
gcc Ala	atg Met	tca Ser	tta Leu 20	cct Pro	ccg Pro	gca Ala	gat Asp	tcc Ser 25	Val	cct Pro	gac Asp	acg Thr	tca Ser 30	tca Ser	cct Pro	96
cca Pro	gct Ala	cct Pro 35	cct Pro	ttg Leu	tct Ser	cct Pro	ctt Leu 40	ccc Pro	cca Pro	cca Pro	ttg Leu	agc Ser 45	tct Ser	cct Pro	ccg Pro	144
ccg Pro	ttg Leu 50	cct Pro	tca Ser	cca Pro	ccg Pro	cct Pro 55	ctc Leu	tcc Ser	gct Ala	ccc Pro	acc Thr 60	gct Ala	tcc Ser	cca Pro	ccg Pro	192
cct Pro 65	ctt Leu	ccg Pro	gtt Val	gaa Glu	tcc Ser 70	cca Pro	ccg Pro	tct Ser	cct Pro	cct Pro 75	ata Ile	gaa Glu	tca Ser	cca Pro	ccg Pro 80	240
cct Pro	cct Pro	cta Leu	ctg Leu	gaa Glu 85	tca Ser	cct Pro	cct Pro	cct Pro	cct Pro 90	ccg Pro	ttg Leu	gaa Glu	tct Ser	cca Pro 95	tct Ser	288
cca Pro	ccg Pro	tct Ser	cct Pro 100	cac His	gtc Val	tca Ser	gct Ala	cct Pro 105	tcc Ser	ggt Gly	tca Ser	ccg Pro	cca Pro 110	tta Leu	ccc Pro	336
ttc Phe	ctt Leu	ccc Pro 115	gcc Ala	aaa Lys	cct Pro	tct Ser	ccg Pro 120	ccg Pro	cct Pro	tct Ser	tca Ser	cct Pro 125	ccc Pro	tcc Ser	gag Glu	384
aca Thr	gtt Val 130	ccg Pro	ccg Pro	gga Gly	aat Asn	acg Thr 135	att Ile	tct Ser	cca Pro	cca Pro	cct Pro 140	cgt Arg	tca Ser	ctt Leu	ccc Pro	432
cc Ser 145	gaa Glu	tca Ser	acc Thr	ccg Prọ	ccg Pro 150	gtg Val	aac Asn	aca Thr	gct Ala	tct Ser 155	cct Pro	cca Pro	ccg Pro	cca Pro	tct Ser 160	480
cct Pro	cct Pro	cgc Arg	cgc Arg	cgt Arg 165	agt Ser	ggc Gly	cct Pro	aag Lys	cct Pro 170	tcg Ser	ttt Phe	cct Pro	cct Pro	ccc Pro 175	atc Ile	528
aat Asn	tct Ser	tct Ser	cca Pro 180	cca Pro	aat Asn	cct Pro	tct Ser	ccg Pro 185	aac Asn	act Thr	ccg Pro	tca Ser	ctc Leu 190	cca Pro	gaa Glu	576
nct Thr	tct Ser	cct Pro 195	cca Pro	cct Pro	aaa Lys	cca Pro	ccg Pro 200	ctc Leu	tca Ser	acg Thr	Thr	cca Pro 205	ttt Phe	ccc Pro	tcc Ser	624

												ctt Leu		672
Phe 225												cct Pro		720
												ata Ile 255		768
												acg Thr		816
			Arg									atc Ile		864
	_				_					_		tac Tyr	_	912
							_	_	_			caa Gln		960
												ggt Gly 335		1008
												aaa Lys		1056
		_		 _	_	_	_				_	gcg Ala		1104
												tca Ser		1152
												ggt Gly		1200
												gcg Ala 415		1248
												caa Gln		1296

----

gag gtt gac aca atc agt agg gtt cat cat aag cac ctc gtt tca ttg Glu Val Asp Thr Ile Ser Arg Val His His Lys His Leu Val Ser Leu gtt ggt tat tgc gtt aat gga gat aaa aga ctc ttg gtt tac gag ttt Val Gly Tyr Cys Val Asn Gly Asp Lys Arg Leu Leu Val Tyr Glu Phe gtt cct aaa gat acc ttg gag ttc cac ttg cat gag aac aga gga agc Val Pro Lys Asp Thr Leu Glu Phe His Leu His Glu Asn Arg Gly Ser gtg ttg gaa tgg gaa atg agg ctc agg att gct gta gga gca gca aaa Val Leu Glu Trp Glu Met Arg Leu Arg Ile Ala Val Gly Ala Ala Lys gga tta gct tat ctt cat gag gat tgc agt cca act ata att cac cgt Gly Leu Ala Tyr Leu His Glu Asp Cys Ser Pro Thr Ile Ile His Arg gat atc aaa gca gct aat atc ctt cta gat tcc aaa ttt gag gca aag Asp Ile Lys Ala Ala Asn Ile Leu Leu Asp Ser Lys Phe Glu Ala Lys gtc tct gac ttt gga cta gcc aag ttt ttc tca gac acc aat tca tca Val Ser Asp Phe Gly Leu Ala Lys Phe Phe Ser Asp Thr Asn Ser Ser ttc act cat atc tct act cga gtg gta gga act ttc gga tac atg gct Phe Thr His Ile Ser Thr Arg Val Val Gly Thr Phe Gly Tyr Met Ala cca gaa tac gcg tcc agt ggt aaa gta act gat aaa tca gat gta tat Pro Glu Tyr Ala Ser Ser Gly Lys Val Thr Asp Lys Ser Asp Val Tyr tcc ttt ggg gtc gtg ctt cta gaa ctc atc act gga cgt cca tca att Ser Phe Gly Val Val Leu Leu Glu Leu Ile Thr Gly Arg Pro Ser Ile ttc gcc aaa gat tct tcc aca aac cag agt tta gta gac tgg gcg agg Phe Ala Lys Asp Ser Ser Thr Asn Gln Ser Leu Val Asp Trp Ala Arg cca ttg ctt acg aaa gca atc tct gga gaa agt ttt gac ttt ctt gta bro Leu Eu Thr Lys Ala Ile Ser Gly Glu Ser Phe Asp Phe Leu Val gac tca agg ttg gag aag aat tac gat aca act cag atg gca aac atg Asp Ser Arg Leu Glu Lys Asn Tyr Asp Thr Thr Gln Met Ala Asn Met 

1	gct Ala	gct Ala	tgt Cys	gct Ala	gct Ala 645	gct Ala	tgc Cys	ata Ile	cgc Arg	caa Gln 650	tca Ser	gct Ala	tgg Trp	ctt Leu	cgg Arg 655	cct Pro	1968
•	aga Arg	atg Met	agc Ser	cag Gln 660	gta Val	gta Val	cgt Arg	gct Ala	ctt Leu 665	gaa Glu	ggc Gly	gag Glu	gtg Val	gcc Ala 670	ctg Leu	aga Arg	2016
	aag Lys	gtc Val	gaa Glu 675	gag Glu	act Thr	Gly 999	aat Asn	agc Ser 680	gtg Val	acc Thr	tat Tyr	agc Ser	tct Ser 685	tct Ser	gaa Glu	aac Asn	2064
	ccg Pro	aat Asn 690	gac Asp	atc Ile	aca Thr	cca Pro	cgg Arg 695	tat Tyr	gga Gly	aca Thr	aat Asn	aag Lys 700	agg Arg	aga Arg	ttc Phe	gac Asp	2112
•	aca Thr 705	ggt Gly	tca Ser	agc Ser	gat Asp	ggt Gly 710	tac Tyr	act Thr	tca Ser	gaa Glu	tat Tyr 715	gga Gly	gtt Val	aac Asn	cct Pro	tct Ser 720	2160
(	cag Gln	tcg Ser	agc Ser	agt Ser	gaa Glu 725	cat His	caa Gln	cag Gln	gtg Val	aat Asn 730	act Thr	tag					2196

<210> 17 <211> 731

<212> PRT

<213> Arabidopsis thaliana

<40	0 > 1	7													
1				5	Ser				10					15	
Ala	Met	Ser	Leu 20	Pro	Pro	Ala	Asp	Ser 25	Val	Pro	Asp	Thr	Ser 30	Ser	Pro
		35			Ser		40					45			
Pro	Leu 50	Pro	Ser	Pro	Pro	Pro 55	Leu	Ser	Ala	Pro	Thr 60	Ala	Ser	Pro	Pro
65					Ser 70					75					80
				85	Ser				90					95	
			100		Val			105					110		
		115			Pro		120					125			
•	130				Asn	135					140	_			
145										155					160
Pro	Pro	Arg	Arg		Ser					Ser		Pro	Pro	Pro 175	Ile
Asn E	Ser	Ser	Pro	Pro	Asn	Pro	Ser	Pro	Asn	Thr	Pro	Ser	Leu	Pro	Glu

Thr Ser Pro Pro Pro Lys Pro Pro Leu Ser Thr Thr Pro Phe Pro Ser Ser Ser Thr Pro Pro Pro Lys Lys Ser Pro Ala Ala Val Thr Leu Pro Phe Phe Gly Pro Ala Gly Gln Leu Pro Asp Gly Thr Val Ala Pro Pro Ile Gly Pro Val Ile Glu Pro Lys Thr Ser Pro Ala Glu Ser Ile Ser Pro Gly Thr Pro Gln Pro Leu Val Pro Lys Ser Leu Pro Val Thr Thr Ser Tyr His Arg Ser Ser Ala Gly Phe Leu Phe Gly Gly Val Ile Val Gly Ala Leu Leu Ile Leu Leu Gly Leu Leu Phe Val Phe Tyr Arg Ala Thr Arg Asn Arg Asn Asn Asn Ser Ser Ser Ala His His Gln Ser Lys Thr Pro Ser Lys Val Gln His His Arg Gly Gly Asn Ala Gly Thr Asn Gln Ala His Val Ile Thr Met Pro Pro Pro Ile His Ala Lys Tyr Ile Ser Ser Gly Gly Cys Asp Thr Lys Glu Asn Asn Ser Val Ala Lys Asn Ile Ser Met Pro Ser Gly Met Phe Ser Tyr Glu Glu Leu Ser Lys Ala Thr Gly Gly Phe Ser Glu Glu Asn Leu Gly Glu Gly Gly Phe Gly Tyr Val His Lys Gly Val Leu Lys Asn Gly Thr Glu Val Ala Val Lys Gln Leu Lys Ile Gly Ser Tyr Gln Gly Glu Arg Glu Phe Gln Ala Glu Val Asp Thr Ile Ser Arg Val His His Lys His Leu Val Ser Leu Val Gly Tyr Cys Val Asn Gly Asp Lys Arg Leu Leu Val Tyr Glu Phe Val Pro Lys Asp Thr Leu Glu Phe His Leu His Glu Asn Arg Gly Ser Val Leu Glu Trp Glu Met Arg Leu Arg Ile Ala Val Gly Ala Ala Lys Gly Leu Ala Tyr Leu His Glu Asp Cys Ser Pro Thr Ile Ile His Arg Asp Ile Lys Ala Ala Asn Ile Leu Leu Asp Ser Lys Phe Glu Ala Lys al Ser Asp Phe Gly Leu Ala Lys Phe Phe Ser Asp Thr Asn Ser Ser Phe Thr His Ile Ser Thr Arg Val Val Gly Thr Phe Gly Tyr Met Ala Pro Glu Tyr Ala Ser Ser Gly Lys Val Thr Asp Lys Ser Asp Val Tyr Ser Phe Gly Val Val Leu Leu Glu Leu Ile Thr Gly Arg Pro Ser Ile Phe Ala Lys Asp Ser Ser Thr Asn Gln Ser Leu Val Asp Trp Ala Arg Pro Leu Leu Thr Lys Ala Ile Ser Gly Glu Ser Phe Asp Phe Leu Val

```
610
                        615
                                             620
Asp Ser Arg Leu Glu Lys Asn Tyr Asp Thr Thr Gln Met Ala Asn Met
                    630
                                         635
Ala Ala Cys Ala Ala Cys Ile Arg Gln Ser Ala Trp Leu Arg Pro
                645
                                     650
Arg Met Ser Gln Val Val Arg Ala Leu Glu Gly Glu Val Ala Leu Arg
                                665
Lys Val Glu Glu Thr Gly Asn Ser Val Thr Tyr Ser Ser Glu Asn
                            680
Pro Asn Asp Ile Thr Pro Arg Tyr Gly Thr Asn Lys Arg Arg Phe Asp
                        695
                                             700
Thr Gly Ser Ser Asp Gly Tyr Thr Ser Glu Tyr Gly Val Asn Pro Ser
                    710
Gln Ser Ser Glu His Gln Gln Val Asn Thr
                725
<210> 18 .
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<220>
<221> misc_feature
<222> (1)..(27)
<223> N(1) =i N(2) =i N(3) =T/c/A N(4) =i N(5) =A/T N(6) =T/C
N(7) = A/g
<400> 18
ggnggtttcg gnatngtntt nnaangg
                                                                  27
<210> 19
<211> 9
<212> PRT
<213> Brassica napus
<220>
<221> PEPTIDE
<222> (1)..(9)
<223> X(1) = F/Y
<400> 19
Gly Gly Phe Gly Ile Val Xaa Lys Gly
<sup>€</sup>210> 20
<211> 22
<212> DNA
<213> Artificial Sequence
```

```
<223> Description of Artificial Sequence: primer
 <220>
 <221> misc feature
 <222> (1)..(22)
 <223> N(1) =i N(2) =i N(3) =T/g N(4) =i N(5) =i N(6) =A/g N(7)
 =A/g N(8) =T/c
 <400> 20
 aanatnenng ceatneenaa nn
                                                                22
 <210> 21
 <211> 8
 <212> PRT
 <213> Brassica napus
 <400> 21
Asp Phe Gly Met Ala Arg Ile Phe
210> 22
<211> 16
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 22
taaccaacaa gagaca
                                                                   16
<210> 23
<211> 23
<212> DNA
<213> Artificial Sequence
₹220>
<223> Description of Artificial Sequence: primer
<220>
<221> misc_feature
<222> (1)..(23)
<223> N(1) =g/A N(2) =i N(3) =g/A N(4) =T/C N(5) =i N(6) =i
                                                                        N(7)
      N(8) = I N(9) = A/g N(10) = A/g
=A/g
<400> 23
annannttng cnannccnaa ntc
                                                          23
```

```
<210> 24
 <211> 8
 <212> PRT
 <213> Arabidopsis thaliana
 <400> 24
 Asp Phe Gly Leu Ala Lys Leu Leu
 <210> 25
 <211> 18
 :212> DNA
 <213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: primer
<400> 25
ccactcccaa ctttcaac
                                                                     18
<210> 26
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
223> Description of Artificial Sequence: primer
<400> 26
ggaaagcttg catgcctgca ggtcgac
                                                                    27
<210> 27
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: primer
<400> 27
cgcctgcagg taatacgact cactataggg
                                                                    30
```